

OM of: US-09-805-550-2 to: Issued\_Patents\_NA:\* out\_format : pfs  
Date: Aug 31, 2002 7:35 AM

About: Results were produced by the GenCore software, version 4.5,  
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## Command line parameters:

-MODEL=frame+p2n.model -DEV=x1h  
-O=/cgn2\_1/USPRO\_pool/US09805550/runat\_29082002.160819.289/app\_query.fasta\_1.897  
-DB=Issued\_Patents\_NA -OPMT=fastap -SUFFIX=oligop2n.rni  
-GAPOP=4.500 -GAPEXT=0.050 -MINMATCH=0.100 -LOOPEXT=0.000  
-IOOPEXT=0.000 -QGAPOP=4.500 -QGAPEXT=0.050 -XGAPOP=60.000  
-XGAPEXT=60.000 -FGAPOP=6.000 -DELEXT=7.000 -YGAPOP=60.000  
-MATRIX=oligo -TRANS=human40.cdi -LIST=45 -DOCALLIGN=200  
-THR\_SCORE=quality -THR\_MIN=1 -ALIGN=15 -MODE=LOCAL -OUTFMT=pfs  
-NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000  
-USER=US9805550.ecgn1\_1.58 -NCPU=6 -ICPU=3 -LONGLOC  
-DEV\_TIMEOUT=120 -WARN\_TIMEOUT=30 -NO\_XLIFY -WAIT -THREADS=1

## Search information block:

Query: US-09-805-550-2  
Query length: 405  
Database: Issued\_Patents\_NA:\*  
Database sequences: 383533  
Database length: 122816752  
Search time (sec): 69.340000

WARN: XGAPOP and YGAPOP must be equal. Assuming YGAPOP=XGAPOP=60.000  
WARN: XGAPEXT and YGAPEXT must be equal. Assuming YGAPEXT=XGAPEXT=60.000

## score\_list:

Sequence	Strd Orig	ZScore	EScore	Len	Documentation	...
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-09-413-574-1 +		405.00	6757.54	0.0	1522	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-09-413-574-3 +		13.00	192.11	0.0052	1702	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-09-103-840A-1 -		9.00	68.43	3.9e+04	4411529	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:5242821-11 +		8.00	119.74	55.57	352	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-822-999-10 -		8.00	114.94	102.88	685	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-791-347-16 -		8.00	114.86	103.99	693	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-869-696-6 +		8.00	113.24	127.93	867	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-869-696-20 +		8.00	111.90	151.92	1044	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-492-027A-5 +		8.00	111.10	168.41	1167	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-869-696-21 +		8.00	110.42	183.71	1282	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-869-696-22 +		8.00	109.93	195.61	1372	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-136-277-18 +		8.00	109.58	204.69	1441	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-479-403-18 +		8.00	109.58	204.69	1441	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-835-734-18 +		8.00	109.58	204.69	1441	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-869-696-8 +		8.00	108.43	237.21	1690	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-714-918-16 -		8.00	107.15	279.51	2018	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-09-265-315-16 -		8.00	107.15	279.51	2018	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-09-265-315-16 -		8.00	107.15	279.51	2018	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-09-266-417-16 -		8.00	107.15	279.51	2018	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-136-277-1 +		8.00	104.91	372.17	2750	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-479-403-1 +		8.00	104.91	372.17	2750	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-835-734-1 +		8.00	104.91	372.17	2750	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-808-982-1 +		8.00	104.25	405.10	3014	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-09-306-902A-1 +		8.00	104.25	405.10	3014	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-693-174-5 +		8.00	103.92	422.85	3157	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-09-253-738-5 +		8.00	103.92	422.85	3157	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-446-038B-2 +		8.00	103.19	464.57	3495	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-446-038B-2 +		8.00	103.19	464.57	3495	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-805-445-2 +		8.00	103.19	464.57	3495	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-064-067D-2 +		8.00	103.19	464.57	3495	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-09-066-208-2 +		8.00	103.19	464.57	3495	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-980-080-3 +		8.00	103.19	464.57	3495	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-026-138E-16 +		8.00	102.27	522.58	3969	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-678-614-1 +		8.00	101.78	556.47	4248	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-231-193A-57 +		8.00	101.06	610.44	4695	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-486-273A-57 +		8.00	101.06	610.44	4695	1
/cgn2_6/prodata/1/ina/6A.COMB.seq:US-08-940-086A-57 +		8.00	101.06	610.44	4695	1
/cgn2_6/prodata/1/ina/6B.COMB.seq:US-08-940-086A-57 +		8.00	101.06	610.44	4695	1

/cgn2\_6/prodata/1/ina/6A.COMB.seq:US-08-869-696-1 + 8.00 99.88 710.12 5529  
/cgn2\_6/prodata/1/ina/6B.COMB.seq:US-08-850-880-11 + 8.00 90.90 2.2e+03 19182  
/cgn2\_6/prodata/1/ina/6B.COMB.seq:US-08-944-916-11 + 8.00 90.90 2.2e+03 19182  
/cgn2\_6/prodata/1/ina/6A.COMB.seq:US-08-846-111D-15 + 8.00 85.54 4.5e+03 4035  
/cgn2\_6/prodata/1/ina/6B.COMB.seq:US-09-103-840A-1 + 8.00 51.68 2.8e+05 44115  
/cgn2\_6/prodata/1/ina/6B.COMB.seq:US-08-751-767A-15 - 7.00 113.96 116.68 77  
/cgn2\_6/prodata/1/ina/6A.COMB.seq:US-08-030-731A-36 - 7.00 113.16 129.25 86

seq\_name: /cgn2\_6/prodata/1/ina/6B.COMB.seq:US-09-413-574-1

## seq\_documentation block:

; Sequence 1, Application US/09413574  
; Patent No. 6235972  
; GENERAL INFORMATION:  
; APPLICANT: Mahajan, Pramod B.  
; APPLICANT: Tagliani, Laura  
; TITLE OF INVENTION: Maize Rad3 Genes and Uses Thereof  
; FILE REFERENCE: 0964  
; CURRENT APPLICATION NUMBER: US/09/413,574  
; EARLIER FILING DATE: 1999-10-06  
; EARLIER APPLICATION NUMBER: 60/109,728  
; EARLIER FILING DATE: 1998-11-23  
; NUMBER OF SEQ ID NOS: 5  
; SOFTWARE: FastSeq for Windows Version 3.0  
; SEQ ID NO 1  
; LENGTH: 1522  
; TYPE: DNA  
; ORGANISM: Zea mays  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (58)...(1272)  
US-09-413-574-1

## alignment\_scores:

Quality: 405.00 Length: 405  
Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

## alignment\_block:

US-09-805-550-2 x US-09-413-574-1 ..  
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1 MetTysLeuAsnValIyThrLeuLysGlyThrsAnpGluIleGluAl 17  
|||||  
58 ATGAAGCTTACGTCATACACCTCAAGGACCAACTTCGATCGAGGC 107  
17 AserProAspAlaSerValAlaAspValLysArgTleIleGluThrThrg 34  
|||||  
108 GAGCCCGATGATCGTGTGATGTGAGAGATCATTTGAGACCACTC 157  
34 InGlyInSerThrTyrArgAlaAspGlnImeTleuIleTyrGlnGly 50  
|||||  
158 AAGGTCAGATGATCTACCGGCGGACCAAGTCTATATACCAAGG 207  
51 LysIleLeuLysAspGluThrThrLeuLysSerAnGlyValAlaGluAs 67  
|||||  
208 AAATTCACAGATGATGATGATGATGATGATGATGATGATGATGATG 257  
84 LaserThraLethrThraLysAlaProAlaThrLeuAlaGlnProAla 100  
|||||  
308 CTTCACCGCTACCTACGCAAAAGCTCTGCAACCTGCGCCCAACCTGCT 357  
101 AlaProValAlaProAlaAlaSerValAlaArgThrProThrGlnAlaPr 117  
|||||  
358 GCCCGTGGGCGCCGCTGATGATGATGATGATGATGATGATGATGATG 407  
117 OvaAlaThrAlaGluThrAlaProProSerValGlnProGlnAlaAlaPr 134

```

|||||
408 TGTGTGCGACAGCTGGAACGGCACCTCCAAAGTGCCAACTCAGGCGTGC 457
134 roAlaAlaThrValAlaAlaThrAspAspAlaAspValTyrSerGlnAla 150
458 CAGCTGCTCAAGGTTGCTGCTACTGATGATGCTGATGCTGATGCTGATG 507
151 AlaSerAsnLeuValPheGlyAsnAsnLeuGluGlnThrTleGlnGln 167
508 GCTTCAAACTGTGATTTGGCAACATCTAGAAACACACTATCCAAACAAT 557
167 eleuAspMetGlyGlyGlyThrTropGluArgAspThrValValArgAla 184
558 TCTTGACATGGGTGGTGTACATGGAGACGTGATCTGTTGTCGTGCTC 607
184 euArGlnAlaAlaTyrAsnAsnProGluArgAlaIleAspTyrLeuTyrSer 200
608 TACGTGCTGCATACATACATACCCGAGAGAGCTATAGCTACCTGCTATCT 657
201 GlyIleProGluAsnValGluAlaGlnProValAlaArgAlaProAlaAl 217
658 GGAATTCCTGAGATGTGGAGGCTCAGCTGTTGCCGAGACCTGCTGC 707
217 aglyGlnGlnThrAsnGlnGlnAlaAlaSerProAlaGlnProAlaVala 234
708 TGGCCACAAACAATCAGACAGCGCATACCCGCTCAGCCAGCAGCGTGG 757
234 IalLeuProValGlnProSerProAlaSerAlaGlyProAsnAlaAsnPro 250
758 CATTGCGCATGCGACGATCAGCTGCTGCGAGGGCTATGCAATTCCT 807
251 LeuAsnLeuPheProGlnGlyValProSerGlyGlySerAsnProGlyVal 267
808 TTGAACCTTTCTCCAGGGTGTTCACAGTGGTGGTCCACCCAGCTGT 857
267 IValProGlyAlaGlySerGlyAlaLeuAspAlaLeuArgGlnLeuProG 284
858 TGTTCAGGTGCGAGATCTGTGCTCTGATGCTTGCACAGCTTCCAC 907
284 IAlPheGlnAlaLeuLeuGlnLeuValAlaAlaAsnProGlnIleLeuGln 300
908 AGTTTCAACACACTCCTTTCAGTTAGTCCAGGCTATCTCAATCTTGCAG 957
301 PrometLeuGlnGlnLeuGlnGlyGlnAsnProGlnIleLeuArgLeu 317
958 CCAATGCTTCAAGAGCTAGTAAACCAACCAATTCCTGGCGTGTGAT 1007
317 eGlnGluAsnGlnAlaGluPheLeuArgLeuValaAsnGluSerProGluG 334
1008 TCAAGGAAATCAAGCTGAGTTCCTCCGCTTGGTGAATGATCTCTGAGG 1057
334 IyGlyProGlyGlyAsnIleLeuGlyGlnLeuAlaAlaValProGln 350
1058 GTGGTCTCGAGGAGACATACTAGTCACTGCGACCTGCTGGCCACAA 1107
351 ThrLeuThrValThrProGluGluArgGluAlaIleGlnArgLeuGlu 367
1108 ACGCTGACAGTTACCCCAAGAGAACGGAGGCTATCCACGCGCTCAGAG 1157
367 yMetGlyPheAsnArgGluLeuValLeuGlnValaPhePheAlaCysAsn 384
1158 AATGGGGTTCAACCGTGAGCTTGTGCTAGAGATTTTCTTTCGATGCCA 1207
384 yAspGlnGluLeuThrAlaAsnTyrLeuLeuAspHisGlyHisGluPhe 400
1208 AGGACAGAACAGGTTACAGCCCACTACCTCTGATCATGGCCATGAGTTT 1257
401 AspAspGlnGlnGln 405
1258 GAGATCAGCAGCA 1272
seq_name: /cgn2_6/ptodata/1/lna/6B_COMB.seq:US-09-413-574-3

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seq_documentation_block:
; Sequence 3, Application US/09413574
; Patent No. 6235972
; GENERAL INFORMATION:
; APPLICANT: Mahajan, Pramod B.
; TITLE OF INVENTION: Maize Rad23 Genes and Uses Thereof
; FILE REFERENCE: 0964
; CURRENT APPLICATION NUMBER: US/09/413,574
; CURRENT FILING DATE: 1999-10-06
; EARLIER APPLICATION NUMBER: 60/109,728
; EARLIER FILING DATE: 1998-11-23
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 3
; LENGTH: 1702
; TYPE: DNA
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (106)...(1209)
US-09-413-574-3

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alignment_scores:
Quality: 13.00 Length: 13
Ratio: 1.000 Gaps: 0
Percent Similarity: 100.000 Percent Identity: 100.000

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alignment_block:
US-09-805-550-2 x US-09-413-574-3 ..

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Align seg 1/1 to: US-09-413-574-3 from: 1 to: 1702

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182 ArgAlaLeuArgAlaAlaTyrAsnAsnProGluArgAla 194
604 AGGCTCTCCGTCGCGCTTACACAAACCCGAGAGCTGCT 642

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seq_name: /cgn2_6/ptodata/1/lna/6B_COMB.seq:US-09-103-840A-1

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seq_documentation_block:
; Sequence 1, Application US/09103840A
; Patent No. 6294328
; GENERAL INFORMATION:
; APPLICANT: FLEISCHMAN, Robert D.
; APPLICANT: WHITE, Owen R.
; APPLICANT: FRASER, Claire M.
; APPLICANT: VENTER, John C.
; TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM
; FILE REFERENCE: TUBERCULOSIS
; FILE REFERENCE: 24366-20007.00
; CURRENT APPLICATION NUMBER: US/09/103,840A
; CURRENT FILING DATE: 1998-06-24
; NUMBER OF SEQ ID NOS: 2
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 4411529
; TYPE: DNA
; ORGANISM: Mycobacterium tuberculosis
; OTHER INFORMATION: H37Rv
US-09-103-840A-1

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alignment_scores:
Quality: 10.00 Length: 10
Ratio: 1.000 Gaps: 0
Percent Similarity: 100.000 Percent Identity: 100.000

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alignment_block:
US-09-805-550-2 x US-09-103-840A-1/rev ..

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Align seg 1/1 to reverse of: US-09-103-840A-1 from: 1 to: 4411529

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99 ProAlaAProValAProAlaAaser 108
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336823 CCGGCGCAGCGGTGCTCCAGCGCATCG 336794

seq_name: /cgn2_6/ptodata/1/ina/backfiles1.seq:5242821-11
seq_documentation_block:
; Patent No. 5242821
; APPLICANT: PALVA, LIKKA, SIBAKHOV, MERVIT
; TITLE OF INVENTION: LACTOCOCCUS PROMOTER AND SIGNAL
; SEQUENCES FOR EXPRESSION IN BACTERIA
; NUMBER OF SEQUENCES: 27
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/377,450
; FILING DATE: 10-JUL-1989
; SEQ ID NO:11:
; LENGTH: 352
5242821-11

alignment_scores:
Quality: 8.00 Length: 8
Ratio: 1.000 Gaps: 0
Percent Similarity: 100.000 Percent Identity: 100.000

alignment_block:
US-09-805-550-2 x 5242821-11 ..
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75 SerlySAlySAaserSer 82
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214 AGTAAAGCCAAAGCCAGTTCATCT 237

seq_name: /cgn2_6/ptodata/1/ina/5B_COMB.seq:US-08-822-999-10
seq_documentation_block:
; Sequence 10, Application US/08822999
; Patent No. 6271026
; GENERAL INFORMATION:
; APPLICANT: Stone, Edwin M.
; APPLICANT: Sheffield, Val C.
; TITLE OF INVENTION: GLAUCOMA COMPOSITIONS AND THERAPEUTIC
; TITLE OF INVENTION: AND DIAGNOSTIC USES THEREFOR
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FOLEY, HOAG & ELIOT LLP
; STREET: One Post Office Square
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02109-2170
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/822,999
; FILING DATE: 21-MAR-1997
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/791,347
; FILING DATE: 30-JAN-1997
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/748,479
; FILING DATE: 08-NOV-1996
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/234,218
; FILING DATE: 28-APR-1994
; ATTORNEY/AGENT INFORMATION:
; NAME: Arnold, Beth E.

REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: UIA-010.27
TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-832-1000
; TELEFAX: 617-832-7000
; INFORMATION FOR SEQ ID NO: 10:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 685 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-822-999-10

alignment_scores:
Quality: 8.00 Length: 8
Ratio: 1.000 Gaps: 0
Percent Similarity: 100.000 Percent Identity: 100.000

alignment_block:
US-09-805-550-2 x US-08-822-999-10/rev ..
Align seg 1/1 to reverse of: US-08-822-999-10 from: 1 to: 685

236 ProValGinProSerProAlaSer 243
|||||
265 CCTGTGACAGCCAGCCAGCTTCC 242

seq_name: /cgn2_6/ptodata/1/ina/5B_COMB.seq:US-08-791-347-16
seq_documentation_block:
; Sequence 16, Application US/08791347
; Patent No. 5885776
; GENERAL INFORMATION:
; APPLICANT: Stone, Edwin M.
; APPLICANT: Sheffield, Val C.
; TITLE OF INVENTION: GLAUCOMA COMPOSITIONS AND THERAPEUTIC
; TITLE OF INVENTION: AND DIAGNOSTIC USES THEREFOR
; NUMBER OF SEQUENCES: 16
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FOLEY, HOAG & ELIOT LLP
; STREET: One Post Office Square
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02109-2170
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/791,347
; FILING DATE: 30-JAN-1997
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Arnold, Beth E.
; REGISTRATION NUMBER: 35,430
; REFERENCE/DOCKET NUMBER: UIA-010.26
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-832-1000
; TELEFAX: 617-832-7000
; INFORMATION FOR SEQ ID NO: 16:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 693 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-791-347-16
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alignment\_scores:  
Quality: 8.00 Length: 8  
Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

## alignment\_block:

US-09-805-550-2 x US-08-791-347-16/rev ..

Align seg 1/1 to reverse of: US-08-791-347-16 from: 1 to: 693

236 ProvaIGlnProserProAlaser 243  
|||||  
265 CCTGTCTCAGCCCGCCCTCTC 242

seq\_name: /cgn2\_6/ptodata/1/ina/6A\_COMB.seq:US-08-869-696-6

## seq\_documentation\_block:

Sequence 6, Application US/08869696C  
Patent No. 6031155  
GENERAL INFORMATION:  
APPLICANT: Cameron-Mills, Verena  
APPLICANT: Lok, Flinn  
APPLICANT: Sinjorgo, Catharina Maria Cornelia  
APPLICANT: Van Den Dool, Ronald Tako Marinus  
APPLICANT: Caspers, Martinus Petrus Maria  
APPLICANT: Van Zeijl-Van Der Valk, Maria Joanna  
TITLE OF INVENTION: ARABINOXYLAN DEGRADATION  
FILE REFERENCE: 11225.01US01  
CURRENT APPLICATION NUMBER: US/08/869,696C  
CURRENT FILING DATE: 1997-06-05  
NUMBER OF SEQ ID NOS: 22  
SOFTWARE: PatentIn Ver. 2.0  
SEQ ID NO 6  
LENGTH: 867  
TYPE: DNA  
ORGANISM: barley  
US-08-869-696-6

## alignment\_scores:

Quality: 8.00 Length: 8  
Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

## alignment\_block:

US-09-805-550-2 x US-08-869-696-6 ..

Align seg 1/1 to: US-08-869-696-6 from: 1 to: 867

80 SerSerSerGlyAlaserThrala 87  
|||||  
534 TCAGTTCAGGGGCTCTACGGCA 557

seq\_name: /cgn2\_6/ptodata/1/ina/6A\_COMB.seq:US-08-869-696-20

## seq\_documentation\_block:

Sequence 20, Application US/08869696C  
Patent No. 6031155  
GENERAL INFORMATION:  
APPLICANT: Cameron-Mills, Verena  
APPLICANT: Lok, Flinn  
APPLICANT: Sinjorgo, Catharina Maria Cornelia  
APPLICANT: Van Den Dool, Ronald Tako Marinus  
APPLICANT: Caspers, Martinus Petrus Maria  
APPLICANT: Van Zeijl-Van Der Valk, Maria Joanna  
TITLE OF INVENTION: ARABINOXYLAN DEGRADATION  
FILE REFERENCE: 11225.01US01  
CURRENT APPLICATION NUMBER: US/08/869,696C  
CURRENT FILING DATE: 1997-06-05  
NUMBER OF SEQ ID NOS: 22  
SOFTWARE: PatentIn Ver. 2.0  
SEQ ID NO 20

LENGTH: 1044  
TYPE: DNA  
ORGANISM: barley  
US-08-869-696-20

## alignment\_scores:

Quality: 8.00 Length: 8  
Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

## alignment\_block:

US-09-805-550-2 x US-08-869-696-20 ..

Align seg 1/1 to: US-08-869-696-20 from: 1 to: 1044

80 SerSerSerGlyAlaserThrala 87  
|||||  
917 TCAGTTCAGGGGCTCTACGGCA 940

seq\_name: /cgn2\_6/ptodata/1/ina/5B\_COMB.seq:US-08-492-027A-5

## seq\_documentation\_block:

Sequence 5, Application US/08492027A  
Patent No. 5912333  
GENERAL INFORMATION:  
APPLICANT: Suzuki, Shoichi  
APPLICANT: Burnell, James N  
TITLE OF INVENTION: DNA Encoding Carbonic Anhydrase  
NUMBER OF SEQUENCES: 9  
CORRESPONDENCE ADDRESS:  
ADDRESSER: Birch, Stewart, Kolasch and Birch  
STREET: P.O. Box 747  
CITY: Falls Church  
STATE: VA  
COUNTRY: USA  
ZIP: 22040-0747  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/492,027A  
FILING DATE:  
CLASSIFICATION: 435  
ATTORNEY/AGENT INFORMATION:  
NAME: Murphy Jr, Gerald M  
REGISTRATION NUMBER: 28,977  
REFERENCE/DOCKET NUMBER: 0760-206  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (703) 205-8000  
TELEFAX: (703) 205-8050  
INFORMATION FOR SEQ ID NO: 5:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1167 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: cDNA to mRNA  
FEATURE:  
NAME/KEY: CDS  
LOCATION: 36..851  
US-08-492-027A-5

## alignment\_scores:

Quality: 8.00 Length: 8  
Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

## alignment\_block:

US-09-805-550-2 x US-08-492-027A-5 ..

Align seg 1/1 to: US-08-492-027A-5 from: 1 to: 1167

100 AAlaIAProValaIAProAlaIa 107  
|||||  
222 GCGGCCCCGTCGCGCCGCGCGC 245

seq\_name: /cgn2\_6/ptodata/1/lna/5a\_COMB.seq:US-08-869-696-21

seq\_documentation\_block:

; Sequence 21, Application US/08869696C  
; Patent No. 6031155  
; GENERAL INFORMATION:  
; APPLICANT: Cameron-Mills, Verena  
; APPLICANT: Lok, Finn  
; APPLICANT: Sinjorgo, Catharina Maria Cornelia  
; APPLICANT: Van Den Dool, Ronald Tako Marinus  
; APPLICANT: Caspers, Martinus Petrus Maria  
; APPLICANT: Van Zeijl-Van Der Valk, Maria Joanna  
; TITLE OF INVENTION: ARABINOXYLAN DEGRADATION  
; FILE REFERENCE: 11225.01US01  
; CURRENT APPLICATION NUMBER: US/08/869,696C  
; CURRENT FILING DATE: 1997-06-05  
; NUMBER OF SEQ ID NOS: 22  
; SOFTWARE: PatentIn Ver. 2.0  
; SEQ ID NO 21  
; LENGTH: 1282  
; TYPE: DNA  
; ORGANISM: barley  
US-08-869-696-21

alignment\_scores:

Quality: 8.00 Length: 8  
Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

alignment\_block:

US-09-805-550-2 x US-08-869-696-21 ..  
Align seg 1/1 to: US-08-869-696-21 from: 1 to: 1282

80 SerSerSergIyAlaSerThraIa 87  
|||||  
1155 TCAAGTTCAGGGGCTCTACGCA 1178

seq\_name: /cgn2\_6/ptodata/1/lna/5a\_COMB.seq:US-08-869-696-22

seq\_documentation\_block:

; Sequence 22, Application US/08869696C  
; Patent No. 6031155  
; GENERAL INFORMATION:  
; APPLICANT: Cameron-Mills, Verena  
; APPLICANT: Lok, Finn  
; APPLICANT: Sinjorgo, Catharina Maria Cornelia  
; APPLICANT: Van Den Dool, Ronald Tako Marinus  
; APPLICANT: Caspers, Martinus Petrus Maria  
; APPLICANT: Van Zeijl-Van Der Valk, Maria Joanna  
; TITLE OF INVENTION: ARABINOXYLAN DEGRADATION  
; FILE REFERENCE: 11225.01US01  
; CURRENT APPLICATION NUMBER: US/08/869,696C  
; CURRENT FILING DATE: 1997-06-05  
; NUMBER OF SEQ ID NOS: 22  
; SOFTWARE: PatentIn Ver. 2.0  
; SEQ ID NO 22  
; LENGTH: 1372  
; TYPE: DNA  
; ORGANISM: barley  
US-08-869-696-22

alignment\_scores:  
Quality: 8.00 Length: 8

Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

alignment\_block:

US-09-805-550-2 x US-08-869-696-22 ..

Align seg 1/1 to: US-08-869-696-22 from: 1 to: 1372

80 SerSerSergIyAlaSerThraIa 87  
|||||  
1245 TCAAGTTCAGGGGCTCTACGCA 1268

seq\_name: /cgn2\_6/ptodata/1/lna/5a\_COMB.seq:US-08-136-277-18

seq\_documentation\_block:

; Sequence 18, Application US/08136277  
; Patent No. 5644045  
; GENERAL INFORMATION:  
; APPLICANT: MANDEL, Jean-Louis  
; APPLICANT: AUBOURG, Patrick  
; APPLICANT: MOSSER, Jean  
; APPLICANT: SARDE, Claude  
; TITLE OF INVENTION: X-LINKED ADRENOLEUKODYSTROPHY GENE AND  
; NUMBER OF SEQUENCES: 23  
; CORRESPONDENCE ADDRESSES:  
; ADDRESSEE: Young & Thompson  
; STREET: 745 South 23rd Street  
; CITY: Arlington  
; STATE: VA  
; COUNTRY: USA  
; ZIP: 22202  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/136,277  
; FILING DATE: 15-OCT-1993  
; CLASSIFICATION: 424  
; ATTORNEY/AGENT INFORMATION:  
; NAME: PATCH, Andrew J.  
; REGISTRATION NUMBER: 32,925  
; REFERENCE/DOCKET NUMBER: B2272  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 703-521-2297  
; TELEFAX: 703-685-0573  
; TELEX: 248425 EMBON  
; INFORMATION FOR SEQ ID NO: 18:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 1441 base pairs  
; TYPE: nucleic acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: DNA (genomic)  
US-08-136-277-18

alignment\_scores:

Quality: 8.00 Length: 8  
Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

alignment\_block:

US-09-805-550-2 x US-08-136-277-18 ..

Align seg 1/1 to: US-08-136-277-18 from: 1 to: 1441

100 AAlaIAProValaIAProAlaIa 107  
|||||  
610 GCGGCTCCTGCTGCTCTGCGGCT 633

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seq_name: /cgn2_6/ptodata/1/ina/5B_COMB.seq:US-08-479-403-18
;
; seq_documentation_block:
; Sequence 18, Application US/08479403
; Patent No. 5869039
; GENERAL INFORMATION:
; APPLICANT: MANDEL, Jean-Louis
; APPLICANT: AUBOURG, Patrick
; APPLICANT: MOSSER, Jean
; APPLICANT: SARDE, Claude
; TITLE OF INVENTION: X-LINKED ADRENOLEUKODYSTROPHY GENE AND
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Young & Thompson
; STREET: 745 South 23rd Street
; CITY: Arlington
; STATE: VA
; COUNTRY: USA
; ZIP: 22202
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/479,403
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: PATCH, Andrew J.
; REGISTRATION NUMBER: 32,925
; REFERENCE/DOCKET NUMBER: B2272DIV
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 703-521-2297
; TELEFAX: 703-685-0573
; TELEX: 248425 EMBON
; INFORMATION FOR SEQ ID NO: 18:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1441 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
; US-08-479-403-18

alignment_scores:
  Quality: 8.00      Length: 8
  Ratio: 1.000      Gaps: 0
  Percent Similarity: 100.000  Percent Identity: 100.000

alignment_block:
US-09-805-550-2 x US-08-479-403-18 ..

Align seg 1/1 to: US-08-479-403-18 from: 1 to: 1441

100 AAlaAlaProvaAlaAlaProAlaAla 107
|||||
610 GCGGCTCTGTGCTGCTGCGGCT 633

seq_name: /cgn2_6/ptodata/1/ina/6A_COMB.seq:US-08-835-734-18
;
; seq_documentation_block:
; Sequence 18, Application US/08835734
; Patent No. 6013769
; GENERAL INFORMATION:
; APPLICANT: MANDEL, Jean-Louis
; APPLICANT: AUBOURG, Patrick
; APPLICANT: MOSSER, Jean
; APPLICANT: SARDE, Claude
; TITLE OF INVENTION: X-LINKED ADRENOLEUKODYSTROPHY GENE AND
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Young & Thompson
; STREET: 745 South 23rd Street
; CITY: Arlington
; STATE: VA
; COUNTRY: USA
; ZIP: 22202
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/835,734
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 536
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/479,403
; FILING DATE: 07-JUN-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: PATCH, Andrew J.
; REGISTRATION NUMBER: 32,925
; REFERENCE/DOCKET NUMBER: B2272DIV
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 703-521-2297
; TELEFAX: 703-685-0573
; TELEX: 248425 EMBON
; INFORMATION FOR SEQ ID NO: 18:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1441 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
; US-08-835-734-18

alignment_scores:
  Quality: 8.00      Length: 8
  Ratio: 1.000      Gaps: 0
  Percent Similarity: 100.000  Percent Identity: 100.000

alignment_block:
US-09-805-550-2 x US-08-835-734-18 ..

Align seg 1/1 to: US-08-835-734-18 from: 1 to: 1441

100 AAlaAlaProvaAlaAlaProAlaAla 107
|||||
610 GCGGCTCTGTGCTGCTGCGGCT 633

seq_name: /cgn2_6/ptodata/1/ina/6A_COMB.seq:US-08-869-696-8
;
; seq_documentation_block:
; Sequence 8, Application US/08869696C
; Patent No. 6031155
; GENERAL INFORMATION:
; APPLICANT: Cameron-Mills, Verena
; APPLICANT: Lok, Flinn
; APPLICANT: Sinjorgo, Catharina Maria Cornelia
; APPLICANT: Van Den Dool, Ronald Tako Marinus
; APPLICANT: Caspers, Martinus Petrus Maria
; APPLICANT: Van Zeijl-Van Der Valk, Maria Joanna
; TITLE OF INVENTION: ARABINOSYLIAN DEGRADATION
; REFERENCE: 11225.01US01
; CURRENT APPLICATION NUMBER: US/08/869,696C
; FILING DATE: 1997-06-05
; NUMBER OF SEQ ID NOS: 22
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO: 8
; LENGTH: 1690
; TYPE: DNA
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! ORGANISM: barley  
US-08-869-696-8

alignment\_scores:  
Quality: 8.00 Length: 8  
Ratio: 1.000 Gaps: 0  
Percent Similarity: 100.000 Percent Identity: 100.000

alignment\_block:  
US-09-805-550-2 x US-08-869-696-8 ..

Align seg 1/1 to: US-08-869-696-8 from: 1 to: 1690

80 SerSerSergIyAlaSerThrAla 87  
|||||  
1364 TCAAGTTCAGGGGCTCTACGCA 1387

